## **RAW SEQUENCE LISTING**

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: 10/381, 2624Source:  $15\omega16$ Date Processed by STIC: 10/31/2006

## ENTERED



IFW16

RAW SEQUENCE LISTING DATE: 10/31/2006
PATENT APPLICATION: US/10/781,262A TIME: 14:10:16

Input Set : A:\Substitute Sequence Listing - 2006-10-25.txt

Output Set: N:\CRF4\10312006\J781262A.raw

3 <110> APPLICANT: Oxxon Therapeutics Limited

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5 <120> TITLE OF INVENTION: RECOMBINANT POXVIRUS
 7 <130> FILE REFERENCE: BWT-PT001.2
 9 <140> CURRENT APPLICATION NUMBER: 10/781,262A
10 <141> CURRENT FILING DATE: 2004-02-18
12 <150> PRIOR APPLICATION NUMBER: US 10/047,761
13 <151> PRIOR FILING DATE: 2002-01-15
15 <150> PRIOR APPLICATION NUMBER: US 09/367,781
16 <151> PRIOR FILING DATE: 1999-11-22
18 <150> PRIOR APPLICATION NUMBER: PCT/GB98/00569
19 <151> PRIOR FILING DATE: 1998-02-23
21 <150> PRIOR APPLICATION NUMBER: GB 9800113.4
22 <151> PRIOR FILING DATE: 1998-01-05
24 <150> PRIOR APPLICATION NUMBER: GB 9703592.7
25 <151> PRIOR FILING DATE: 1997-02-21
27 <160> NUMBER OF SEQ ID NOS: 3
29 <170> SOFTWARE: PatentIn version 3.3
31 <210> SEQ ID NO: 1
32 <211> LENGTH: 897
33 <212> TYPE: DNA
34 <213> ORGANISM: Vaccinia virus
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                                                                         120
41 atgtactcgt tagtatttgt tattttgatg tgtataccat ttagttttca aacagtgtat
                                                                         180
43 qatqataaat cggtatgcga ttctgacaat aaagaatata tgggaataga agtttatgta
                                                                         240
45 qaaqcaacgc tagacgaacc cctcagacaa acaacgtgtg aatccaaaat ccataaatat
                                                                         300
47 qqtqcatctq tatcaaacqq aggattaaat atttctqttq atctattaaa ctqttttctt
                                                                         360
49 aattttcata cagttggtgt atacactaat cgcgataccg tatacgcgaa gtttgctagt
                                                                         420
51 ttggatccat ggactacgga acctataaat tctatgaccc atgacgatct agtaaaatta
                                                                         480
53 acagaagaat gtatagtgga catttattta aaatgtgaag tggataaaac aaaggatttc
                                                                         540
55 atgaaaacta acggtaatag attaaaacca agagacttta aaactgttcc tccttctaat
                                                                         600
57 gtaggaagca tgatagaact acagtetgac tattgcgtaa acgatgtgac tacatacgte
                                                                         660
59 aaaatatacg atgagtgtgg aaacattaaa cagcattcca ttccaacact aagagattat
                                                                         720
61 tttaccacca agaatggtca accacgtaaa atattaaaga aaaaatttga taattgttaa
                                                                         780
63 ttgttatttt tataaaaaca agaacggtac ggcgatattt atttttttct aaaacatcta
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69 <211> LENGTH: 219
70 <212> TYPE: PRT
71 <213> ORGANISM: Vaccinia virus
73 <400> SEQUENCE: 2
75 Met Tyr Ser Leu Val Phe Val Ile Leu Met Cys Ile Pro Phe Ser Phe
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76	l				5					10					15	
		Chr	Val	Tvr	_	Asp	Lvs	Ser	Val		Asp	Ser	Asp	Asn	Lvs	Glu
80				20		<u>F</u>	-1-		25	- 4	•			30	•	
	Cyr N	/let	Gly	Ile	Glu	Val	Tyr	Val	Glu	Ala	Thr	Leu	Asp	Glu	Pro	Leu
84	•		35				-	40					45			
87 2	Arg (	Gln	Thr	Thr	Cys	Glu	Ser	Lys	Ile	His	Lys	Tyr	Gly	Ala	Ser	Val
88		50			-		55	-			_	60	_			
91 8	Ser A	Asn	Gly	Gly	Leu	Asņ	Ile	Ser	Val	Asp	Leu	Leu	Asn	Cys	Phe	Leu
92 6	55					70					75					80
95 A	Asn I	Phe	His	Thr	Val	Gly	Val	Tyr	Thr	Asn	Arg	Asp	Thr	Val	Tyr	Ala
96					85					90					95	
99 1	Lys I	Phe	Ala	Ser	Leu	Asp	Pro	$\mathtt{Trp}$	Thr	Thr	Glu	Pro	Ile	Asn	Ser	Met
100				100					10					110		
103	Thr	His	Asp	As <u>r</u>	Let	ı Val	. Lys	s Le	a Thi	r Gli	ı Glı	ı Cys	s Ile	e Val	l Ası	Ile
104			115					12					129			
107	Tyr			S Cys	s Gli	ı Val			s Th:	r Lys	s Asp			Lys	Thi	Asn
108	_	130					139			_		140		_	_	_
	_	Asn	Arg	J Lei	ı Lys			g As	p Phe	e Lys			l Pro	Pro	sei	Asn
	145	~-7	_		7	150		<b>~</b> 1			159	-			. 7	160
	Val	GLY	Sei	r Met			ı Lei	1 GI	n se:	r Ası 170		c Cy	s va.	L ASI		val
116	mla sa	mb -	. П	. 375]	165		. П	~ 7~	~ C1,		_	. 70	. T]	. Tar	179	
	THE	THE	Tyl			3 116	: 1y	L AS	18!		s GI	ASI	.1 116	: шу: 190		n His
120	C0.x	т1.	D~c	180 Th:		1 7 20	, 7\c1	о Пат			r Thi	c T.37	e Δeι			n Pro
124	261	116	199		. ше	ı Arç	, vol	20		_ 1111		. <u>.</u> .	20!		, 011	
	Ara	Lvs			ı Tıvs	s Lys	LVS			o Asi	ı Cvs	3	20.			
128	**** 9	210			,.	, _,.	21!		J							
	<210			D NO	): 3						,					
				ΓH: 2												
133	<212	2> I	YPE	PR'	ľ											
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139					5					10					15	
	Met	Pro	Ala		: Le	ı Glr	ı Glı	n Se		r Sei	r Se	r Se:	r Se		r Cy:	s Thr
143	_	_		20	_	•	•		25		_			30	_	
	Glu	Glu		ı Ası	ı Lya	s His	s His		t GI	A IT	e As	o Va		e 116	э гу	s Val
147	1		35			. m)	. 5	40			<b></b>	. T1	45	- 01-		
			GII	ı Ası	) GII	n Thi		o Tn	r As	n Ası	о га			3 G11	ı se	r Val
151		50	т1.	mh.			55		~ 7 ~	n Dw	- A a ı	60		, 170		, Cer
		GIU	1 116	= THI	. GII	1 Se1 70	. GII	ı se	L AS	o Pro	75 AS	יים י	) GT	ı va.	r GT/	Ser 80
155		Λar	. 7\ ~ 7		c ሞክ፣		c Va	1 G1:	11 7\c1	o 175.		n Dro	o Pro	ን ሞክነ	r Thi	r Tyr
158	GIU	ASL	, wal	ر عدد ر	85	. <i>s</i> e1	. va.	. G1	u AS	90	- vol	, F1(	J FI	- 111	95	y-
	ጥኒታዮ	Ser	· Tl2	ъ Т1 a		, Gls	, G1v	v T.e.	ιι Αν		. Agı	n Ph	e Gla	v Ph		r Lys
163	- y -	DET		100		, 5-1	. 01	L	10		1.01			110		
	Cvs	Pro	Glr			s Sei	: Ile	e Se			r Ala	a Ası	o Gl			r Val
167	-1.5		115					12					12			
	Asn	Ala			ı Sei	r Sei	va:			o Gl	y Gl	ı Gl	у Ьу	s Ası	Se:	r Pro
			-	-						-	-		-	-		

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175	145					150					155					160
178	Ser	Ile	Asp	Ile	Arg	Cys	Ser	Glu	Glu	Glu	Lys	Asp	Ser	Asp	Ile	Lys
179					165					170					175	
182	Thr	His	${\tt Pro}$	Val	Leu	Gly	Ser	Asn	Ile	Ser	His	Lys	Lys	Val	Ser	Tyr
183				180					185					190		
186	Glu	Asp	Ile	Ile	Gly	Ser	Thr	Ile	Val	Asp	Thr	Lys	Cys	Val	Lys	Asn
187			195					200					205			•
190	Leu	Glu	Phe	Ser	Val	Arg	Ile	Gly	Asp	Met	Cys	Lys	Glu	Ser	Ser	Glu
191		210					215					220				
194	Leu	Glu	Val	Lys	Asp	Gly	Phe	Lys	Tyr	Val	Asp	Gly	Ser	Ala	Ser	
	225					230					235					240
198	Gly	Ala	Thr	Asp	Asp	Thr	Ser	Leu	Ile	Asp	Ser	Thr	Lys	Leu		Ala
199					245					250					255	
202	Cys	Val														

VERIFICATION SUMMARY

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